

aaa gat cca agt gga tta tct gga gat gtt ctg ata aat gga gca ccg 336
Lys Asp Pro Ser Gly Leu Ser Gly Asp Val Leu Ile Asn Gly Ala Pro

100	105	110	
cga cct gcc aat ttc aaa tgt aat tca ggt tac gtg gta caa gat gat			384
Arg Pro Ala Asn Phe Lys Cys Asn Ser Gly Tyr Val Val Gln Asp Asp			
115	120	125	
gtt gtg atg ggc act ctg acg gtg aga gaa aac tta cag ttc tca gca			432
Val Val Met Gly Thr Leu Thr Val Arg Glu Asn Leu Gln Phe Ser Ala			
130	135	140	
gct ctt cgg ctt gca aca act atg acg aat cat gaa aaa aac gaa cgg			480
Ala Leu Arg Leu Ala Thr Thr Met Thr Asn His Glu Lys Asn Glu Arg			
145	150	155	160
att aac agg gtc att caa gag tta ggt ctg gat aaa gtg gca gac tcc			528
Ile Asn Arg Val Ile Gln Glu Leu Gly Leu Asp Lys Val Ala Asp Ser			
165	170	175	
aag gtt gga act cag ttt atc cgt ggt gtg tct gga gga gaa aga aaa			576
Lys Val Gly Thr Gln Phe Ile Arg Gly Val Ser Gly Gly Glu Arg Lys			
180	185	190	
agg act agt ata gga atg gag ctt atc act gat cct tcc atc ttg ttc			624
Arg Thr Ser Ile Gly Met Glu Leu Ile Thr Asp Pro Ser Ile Leu Phe			
195	200	205	
ttg gat gag cct aca act ggc tta gac tca agc aca gca aat gct gtc			672
Leu Asp Glu Pro Thr Thr Gly Leu Asp Ser Ser Thr Ala Asn Ala Val			
210	215	220	
ctt ttg ctc ctg aaa agg atg tct aag cag gga cga aca atc atc ttc			720
Leu Leu Leu Leu Lys Arg Met Ser Lys Gln Gly Arg Thr Ile Ile Phe			
225	230	235	240
tcc att cat cag cct cga tat tcc atc ttc aag ttg ttt gat agc ctc			768
Ser Ile His Gln Pro Arg Tyr Ser Ile Phe Lys Leu Phe Asp Ser Leu			
245	250	255	
acc tta ttg gcc tca gga aga ctt atg ttc cac ggg cct gct cag gag			816
Thr Leu Leu Ala Ser Gly Arg Leu Met Phe His Gly Pro Ala Gln Glu			
260	265	270	
gcc ttg gga tac ttt gaa tca gct ggt tat cac tgt gag gcc tat aat			864
Ala Leu Gly Tyr Phe Glu Ser Ala Gly Tyr His Cys Glu Ala Tyr Asn			
275	280	285	
aac cct gca gac ttc ttc ttg gac atc att aat gga gat tcc act gct			912
Asn Pro Ala Asp Phe Phe Leu Asp Ile Ile Asn Gly Asp Ser Thr Ala			
290	295	300	
gtg gca tta aac aga gaa gaa gac ttt aaa gcc aca gag atc ata gag			960
Val Ala Leu Asn Arg Glu Glu Asp Phe Lys Ala Thr Glu Ile Ile Glu			

305	310	315	320	
cct tcc aag cag gat aag cca ctc ata gaa aaa tta gcg gag att tat				1008
Pro Ser Lys Gln Asp Lys Pro Leu Ile Glu Lys Leu Ala Glu Ile Tyr				
	325	330	335	
gtc aac tcc tcc ttc tac aaa gag aca aaa gct gaa tta cat caa ctt				1056
Val Asn Ser Ser Phe Tyr Lys Glu Thr Lys Ala Glu Leu His Gln Leu				
	340	345	350	
tcc ggg ggt gag aag aag aag aag atc aca gtc ttc aag gag atc agc				1104
Ser Gly Gly Glu Lys Lys Lys Lys Ile Thr Val Phe Lys Glu Ile Ser				
	355	360	365	
tac acc acc tcc ttc tgt cat caa ctc aga tgg gtt tcc aag cgt tca				1152
Tyr Thr Thr Ser Phe Cys His Gln Leu Arg Trp Val Ser Lys Arg Ser				
	370	375	380	
ttc aaa aac ttg ctg ggt aat ccc cag gcc tct ata gct cag atc att				1200
Phe Lys Asn Leu Leu Gly Asn Pro Gln Ala Ser Ile Ala Gln Ile Ile				
	385	390	395	400
gtc aca gtc gta ctg gga ctg gtt ata ggt gcc att tac ttt ggg cta				1248
Val Thr Val Val Leu Gly Leu Val Ile Gly Ala Ile Tyr Phe Gly Leu				
	405	410	415	
aaa aat gat tct act gga atc cag aac aga gct ggg gtt ctc ttc ttc				1296
Lys Asn Asp Ser Thr Gly Ile Gln Asn Arg Ala Gly Val Leu Phe Phe				
	420	425	430	
ctg acg acc aac cag tgt ttc agc agt gtt tca gcc gtg gaa ctc ttt				1344
Leu Thr Thr Asn Gln Cys Phe Ser Ser Val Ser Ala Val Glu Leu Phe				
	435	440	445	
gtg gta gag aag aag ctc ttc ata cat gaa tac atc agc gga tac tac				1392
Val Val Glu Lys Lys Leu Phe Ile His Glu Tyr Ile Ser Gly Tyr Tyr				
	450	455	460	
aga gtg tca tct tat ttc ctt gga aaa ctg tta tct gat tta tta ccc				1440
Arg Val Ser Ser Tyr Phe Leu Gly Lys Leu Leu Ser Asp Leu Leu Pro				
	465	470	475	480
atg agg atg tta cca agt att ata ttt acc tgt ata gtg tac ttc atg				1488
Met Arg Met Leu Pro Ser Ile Ile Phe Thr Cys Ile Val Tyr Phe Met				
	485	490	495	
tta gga ttg aag cca aag gca gat gcc ttc ttc gtt atg atg ttt acc				1536
Leu Gly Leu Lys Pro Lys Ala Asp Ala Phe Phe Val Met Met Phe Thr				
	500	505	510	
ctt atg atg gtg gct tat tca gcc agt tcc atg gca ctg gcc ata gca				1584
Leu Met Met Val Ala Tyr Ser Ala Ser Ser Met Ala Leu Ala Ile Ala				

Lys Ser Gly Phe Leu Pro Cys Arg Lys Pro Val Glu Lys Glu Ile Leu
50 55 60

Ser Asn Ile Asn Gly Ile Met Lys Pro Gly Leu Asn Ala Ile Leu Gly
65 70 75 80

Pro Thr Gly Gly Gly Lys Ser Ser Leu Leu Asp Val Leu Ala Ala Arg
85 90 95

Lys Asp Pro Ser Gly Leu Ser Gly Asp Val Leu Ile Asn Gly Ala Pro
100 105 110

Arg Pro Ala Asn Phe Lys Cys Asn Ser Gly Tyr Val Val Gln Asp Asp
115 120 125

Val Val Met Gly Thr Leu Thr Val Arg Glu Asn Leu Gln Phe Ser Ala
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Ala Leu Arg Leu Ala Thr Thr Met Thr Asn His Glu Lys Asn Glu Arg
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Ile Asn Arg Val Ile Gln Glu Leu Gly Leu Asp Lys Val Ala Asp Ser
165 170 175

Lys Val Gly Thr Gln Phe Ile Arg Gly Val Ser Gly Gly Glu Arg Lys
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195 200 205

Leu Asp Glu Pro Thr Thr Gly Leu Asp Ser Ser Thr Ala Asn Ala Val
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Leu Leu Leu Leu Lys Arg Met Ser Lys Gln Gly Arg Thr Ile Ile Phe
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Ser Ile His Gln Pro Arg Tyr Ser Ile Phe Lys Leu Phe Asp Ser Leu
245 250 255

Thr Leu Leu Ala Ser Gly Arg Leu Met Phe His Gly Pro Ala Gln Glu
260 265 270

Ala Leu Gly Tyr Phe Glu Ser Ala Gly Tyr His Cys Glu Ala Tyr Asn
275 280 285

Asn Pro Ala Asp Phe Phe Leu Asp Ile Ile Asn Gly Asp Ser Thr Ala
290 295 300

Val Ala Leu Asn Arg Glu Glu Asp Phe Lys Ala Thr Glu Ile Ile Glu
305 310 315 320

Pro Ser Lys Gln Asp Lys Pro Leu Ile Glu Lys Leu Ala Glu Ile Tyr
325 330 335

Val Asn Ser Ser Phe Tyr Lys Glu Thr Lys Ala Glu Leu His Gln Leu
340 345 350

Ser Gly Gly Glu Lys Lys Lys Lys Ile Thr Val Phe Lys Glu Ile Ser
355 360 365

Tyr Thr Thr Ser Phe Cys His Gln Leu Arg Trp Val Ser Lys Arg Ser
370 375 380

Phe Lys Asn Leu Leu Gly Asn Pro Gln Ala Ser Ile Ala Gln Ile Ile
385 390 395 400

Val Thr Val Val Leu Gly Leu Val Ile Gly Ala Ile Tyr Phe Gly Leu
405 410 415

Lys Asn Asp Ser Thr Gly Ile Gln Asn Arg Ala Gly Val Leu Phe Phe
420 425 430

Leu Thr Thr Asn Gln Cys Phe Ser Ser Val Ser Ala Val Glu Leu Phe
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Val Val Glu Lys Lys Leu Phe Ile His Glu Tyr Ile Ser Gly Tyr Tyr
450 455 460

Arg Val Ser Ser Tyr Phe Leu Gly Lys Leu Leu Ser Asp Leu Leu Pro
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Met Arg Met Leu Pro Ser Ile Ile Phe Thr Cys Ile Val Tyr Phe Met
 485 490 495

Leu Gly Leu Lys Pro Lys Ala Asp Ala Phe Phe Val Met Met Phe Thr
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Leu Met Met Val Ala Tyr Ser Ala Ser Ser Met Ala Leu Ala Ile Ala
 515 520 525

Ala Gly Gln Ser Val Val Ser Val Ala Thr Leu Leu Met Thr Ile Cys
 530 535 540

Phe Val Phe Met Met Ile Phe Ser Gly Leu Leu Val Asn Leu Thr Thr
 545 550 555 560

Ile Ala Ser Trp Leu Ser Trp Leu Gln Tyr Phe Ser Ile Pro Arg Tyr
 565 570 575

Gly Phe Thr Ala Leu Gln His Asn Glu Phe Leu Gly Gln Asn Phe Cys
 580 585 590

Pro Gly Leu Asn Ala Thr Gly Asn Asn Pro Cys Asn Tyr Ala Thr Cys
 595 600 605

Thr Gly Glu Glu Tyr Leu Val Lys Gln Gly Ile Asp Leu Ser Pro Trp
 610 615 620

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